

Message

From: GeneConvene Virtual Institute [contact@geneconvene.org]
Sent: 3/22/2022 5:09:00 AM
To: Kirk, Cassandra [kirk.cassandra@epa.gov]
Subject: Nice review article on Wolbachia and Gene Drive approaches to vector control; Modeling gene drives in Africa

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03/22/2022



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Sharing knowledge to facilitate understanding of gene drive and other genetic biocontrol technologies

Gene Drive & Genetic Biocontrol Newsletter

March 22, 2022

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Genetic biocontrol, also known as genetic pest management and genetic control, is an active research space. Here we have aggregated research and media reports that have appeared since the previous Newsletter.

GeneConvene Global Collaborative Webinars

Registration



NOTE: Atypical Time!!

THURSDAY March 24, 2022

11:00-12:30 EST

Daven Presgraves, Ph.D.

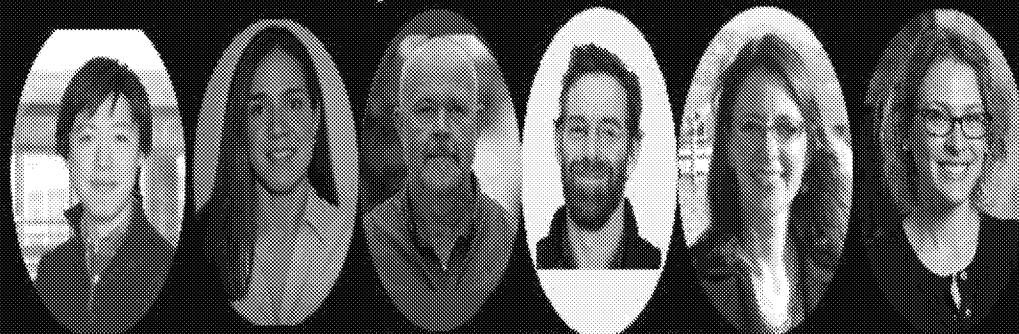
University of Rochester

***Selfish genes, sex
chromosomes, and
speciation in
Drosophila***



Genetic Drive Systems in Nature

A Webinar Series by the GeneConvene Global Collaborative



11:00 am- 12:30pm EST

- March 2 Takashi Akera, Ph.D. Cell biology of female meiotic drive in mice
- March 9 Amanda Larracuenta, Ph.D. Meiotic drive and satellite DNA in *Drosophila*
- March 16 Andreas Houben, Ph.D. Supernumerary B chromosomes : Champions of chromosome drive
- March 24 Daven Presgraves, Ph.D. Selfish genes, sex chromosomes, and speciation in *Drosophila*
- March 30 Anna Lindholm, Ph.D. Ecology of the t-haplotype: A natural gene drive system in house mice
- April 6 Sarah Zanders, Ph.D. WTF spore killers

register here: fnih.org/GeneticDriveSystems

Registration

New Video

Upcoming Meeting!

Gordon Research Conference

The power of gene editing

The Economist

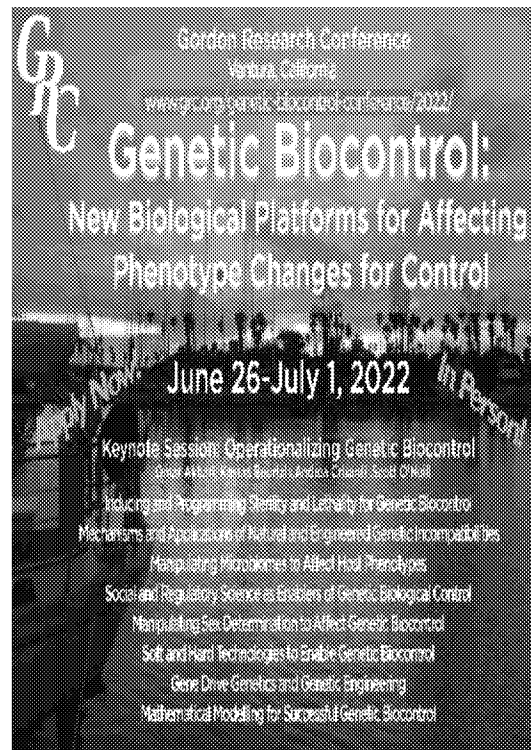
Technologies such as genetic modification and 'CRISPR' will cure hereditary diseases, produce disease-resistant crops and enable the breeding of malaria-free mosquitos. But advances bring ethical and practical dilemmas. Genetically modified food is banned in the EU, and doctors worry that screening for genetic diseases may pave the way for more controversial uses, such as creating so-called designer babies. This film looks at the risks and rewards of gene editing. Gene drive technologies are considered



GENETIC BIOCONTROL

June 26-July 1, 2022

In-Person; Ventura, California



*A convening of world-experts
to share the latest results, technologies,
and best practices.*

New Research and Scholarship

Review Article

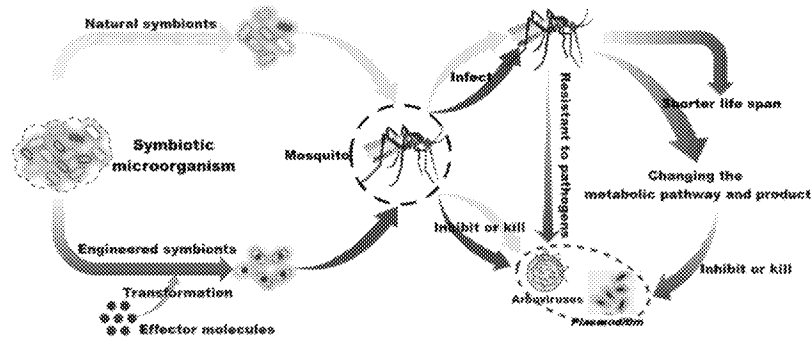
3/18/2021

Symbionts and gene drive: two strategies to combat vector-borne disease

Wang, G.-H., Du, J., Chu, C. Y., Madhav, M., Hughes, G. L. and Champer, J.

Trends in Genetics

<https://doi.org/10.1016/j.tig.2022.02.013>



Mosquitoes bring global health problems by transmitting parasites and viruses such as malaria and dengue. Unfortunately, current insecticide-based control strategies are only moderately effective because of high cost and resistance. Thus, scalable, sustainable, and cost-effective strategies are needed for mosquito-borne disease control. Symbiont-based and genome engineering-based approaches provide new tools that show promise for meeting these criteria, enabling modification or suppression approaches. Symbiotic bacteria like *Wolbachia* are maternally inherited and manipulate mosquito host reproduction to enhance their vertical transmission. Genome engineering-based gene drive methods, in which mosquitoes are genetically altered to spread drive alleles throughout wild populations, are also proving to be a potentially powerful approach in the laboratory. Here, we review the latest developments in both symbionts and gene drive-based methods. We describe some notable similarities, as well as distinctions and obstacles, relating to these promising technologies.

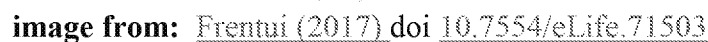
Ecological Impacts of Genetic Biocontrol

3/16/2022 Could species-focused suppression of *Aedes aegypti*, the yellow fever mosquito, and *Aedes albopictus*, the tiger mosquito, affect interacting predators? An evidence synthesis from the literature

Bonds, J. A. S., Collins, C. M. and Gouagna, L.-C.

Pest Management Science

<https://doi.org/10.1002/ps.6870>



2/15/2022 Adult mosquito predation and potential impact on the sterile insect technique

Bimbilé Somda, N. S., Maïga, H., Mamai, W., Bakhoun, T., Wallner, T., Poda, S. B.,
et al.

Scientific Reports

<https://doi.org/10.1038/s41598-022-06565-1>

The sterile insect technique is a promising environmentally friendly method for mosquito control. This technique involves releasing laboratory-produced sterile males into a target field site, and its effectiveness may be affected by the extent of adult mosquito predation. Sterile males undergo several treatments. Therefore, it is vital to understand which treatments are essential in minimizing risks to predation once released. The present study investigates the predation propensity of four mantis species (*Phyllocrania paradoxa*, *Hymenopus coronatus*, *Blepharopsis mendica*, *Deroplatys desiccata*) and two gecko species (*Phelsuma standingi*, *P. laticauda*) on adult *Aedes aegypti*, *Ae. albopictus* and *Anopheles arabiensis* mosquitoes in a laboratory setting. First, any inherent predation preferences regarding mosquito species and sex were evaluated. Subsequently, the effects of chilling, marking, and irradiation, on predation rates were assessed. The selected predators effectively

Wolbachia Anti-Pathogen Effect

3/16/2022

Differential viral RNA methylation contributes to pathogen blocking in
Wolbachia-colonized arthropods

Bhattacharya, T., Yan, L., Crawford, J. M., Zaher, H., Newton, I. L. G. and
Hardy, R. W.

PLoS Pathog

<https://doi.org/10.1371/journal.ppat.1010393>

Arthropod endosymbiont *Wolbachia pipientis* is part of a global biocontrol strategy to reduce the replication of mosquito-borne RNA viruses such as alphaviruses. We previously demonstrated the importance of a host cytosine methyltransferase, DNMT2, in *Drosophila* and viral RNA as a cellular target during pathogen-blocking. Here we report a role for DNMT2 in *Wolbachia*-induced alphavirus inhibition in *Aedes* species. Expression of DNMT2 in mosquito tissues, including the salivary glands, is elevated upon virus infection. Notably, this is suppressed in *Wolbachia*-colonized animals, coincident with reduced virus replication and decreased infectivity of progeny virus. Ectopic expression of DNMT2 in cultured *Aedes* cells is proviral, increasing progeny virus infectivity, and this effect of DNMT2 on virus replication and infectivity is dependent on its methyltransferase activity. Finally, examining the effects of *Wolbachia* on modifications of viral RNA by LC-MS show a decrease in the amount of 5-methylcytosine modification consistent with the down-regulation of DNMT2 in *Wolbachia* colonized mosquito cells and animals. Collectively, our findings support the conclusion that disruption of 5-methylcytosine modification of viral RNA is a vital mechanism operative in pathogen blocking. These data also emphasize the essential role of epitranscriptomic modifications in regulating

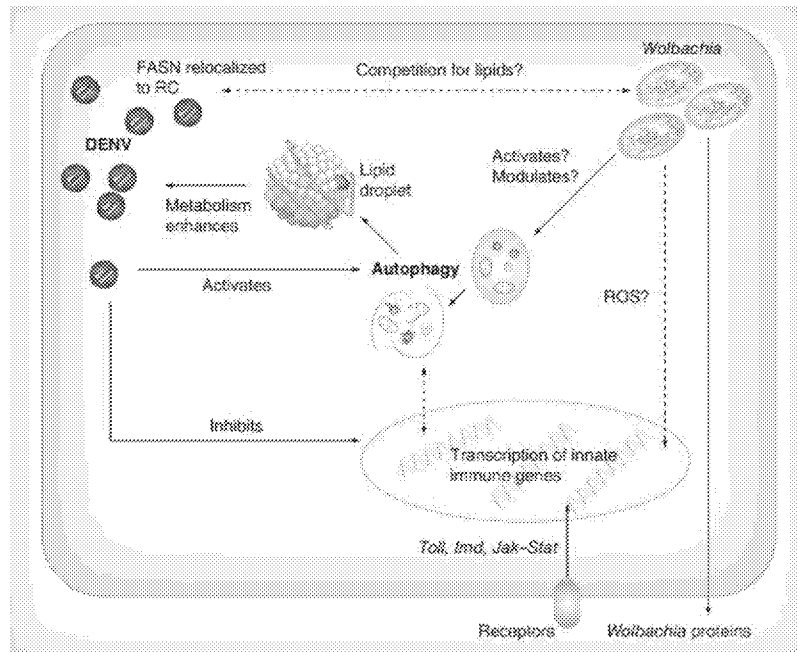


image from: Sinkins (2013) doi.10.2217/fmb.13.95

Modeling Gene Drives

3/18/2022

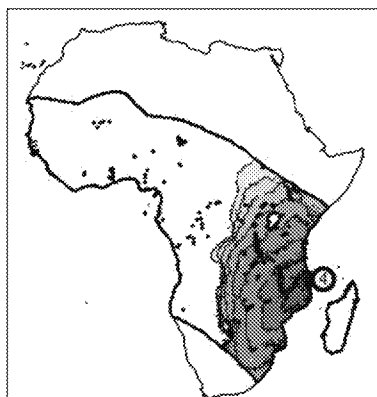
Spatial modelling for population replacement of mosquito vectors at continental scale

Beeton, N. J., Wilkins, A., Ickowicz, A., Hayes, K. R. and Hosack, G. R.

bioRxiv

<https://doi.org/10.1101/2021.10.06.463299>

Malaria is one of the deadliest vector-borne diseases in the world. Researchers are developing new genetic and conventional vector control strategies to attempt to limit its burden. Novel control strategies require detailed safety assessment to ensure responsible and successful deployments.



Anopheles gambiae sensu stricto (s.s.) and

Anopheles coluzzii, two closely related subspecies within the species complex Anopheles gambiae sensu lato (s.l.), are among the dominant malaria vectors in sub-Saharan Africa. These two subspecies readily hybridise and compete in the wild and are also known to have distinct niches, each with spatially and temporally varying carrying capacities driven by precipitation and land use factors. We model the spread and persistence of a population-modifying gene drive system in these subspecies across sub-Saharan Africa by simulating introductions of genetically modified mosquitoes across the African

3/15/2022

Modelling homing suppression gene drive in haplodiploid organisms

Liu, Y. and Champer, J.

bioRxiv

<https://doi.org/10.1101/2021.10.12.464047>

Gene drives have shown great promise for suppression of pest populations. These engineered alleles can function by a variety of mechanisms, but the most common is the CRISPR homing drive, which converts wild-type alleles to drive alleles in the germline of heterozygotes. Some potential target species are haplodiploid, in which males develop from unfertilized eggs and thus have only one copy of each chromosome. This prevents drive conversion, a substantial disadvantage compared to diploids where drive conversion can take place in both sexes. Here, we study homing suppression gene drives in haplodiploids and find that a drive targeting a female fertility gene could still be successful. However, such drives are less powerful than in diploids and suffer more from functional resistance alleles. They are substantially more vulnerable to high resistance

More Research and Scholarship in the Virtual Institute

New Media Coverage

3/19/2022

Squashing malaria could save as many lives as covid-19 has taken

Anonymous

The Economist

<https://www.economist.com/graphic-detail/2022/03/19/squashing-malaria-could-save-as-many-lives-as-covid-19-has-taken>

When it comes to covid-19 vaccines, poor countries in Africa have been stuck at the back of the queue. However, the continent's long wait for another immunological miracle appears to be drawing to a close. Later this year, the world's

3/16/2022

[Studying the active role of the maize B chromosome in the modulation of gene expression](#)

University of Missouri

Phys Org

<https://phys.org/news/2022-03-role-maize-chromosome-modulation-gene.html>

A team of University of Missouri biologists has made a new discovery that provides novel insights into the function and properties of the maize B chromosome. The study was led by Dr. Xiaowen Shi and Dr. Hua Yang, postdoctoral fellows

3/16/2022

[Billions of GE Mosquitoes May Soon Be Released in California and Florida](#)

Mitra, A. N.

Earth Island Journal

<https://www.earthisland.org/journal/index.php/articles/entry/billions-of-ge-mosquitoes-may-soon-be-released-in-california-and-florida/>

The US Environmental Protection Agency's decision last week to allow the release of billions of genetically engineered mosquitoes in California and Florida has several environmental and public health groups worried about the potential impacts of the experimental releases on public

3/15/2022

[Fall armyworms with offspring-killing gene tested on farms in Brazil](#)

Le Page, M.

New Scientist

<https://www.newscientist.com/article/2312341-fall-armyworms-with-offspring-killing-gene-tested-on-farms-in-brazil/>

Fall armyworms genetically modified to wipe out wild populations of the pests have been released in corn fields in São Paulo State in Brazil in the first farm trial of the new technology. The test was a success and is

3/15/2022

[California's first lab-grown mosquitoes may take flight—stirring controversy](#)

Krieger, L. M.

Phys Org

<https://phys.org/news/2022-03-california-lab-grown-mosquitoes-flightstirring-controversy.html>

A biotech firm is seeking permission to release genetically modified mosquitoes into the open air of California for the first time later this year, aiming to reduce the expanding populations of invasive mosquitoes and prevent deadly disease. The controversial research

3/14/2022

[Genetically modified mosquitoes kill their own offspring](#)

Ward, C.

SYFY

<https://www.syfy.com/syfy-wire/genetic-modification-killing-mosquitos-before-theyre-born>

Oxitec, a biotech company, has developed a strain of *A. aegypti* which they call Friendly. Their modified mosquitos are all males and carry a gene which prevents the birth of females in subsequent generations. In short, they're reducing disease-carrying mosquitos

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